



Gut Clone (OTU)

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Figure S1: Collector's curves. The horizontal brackets in each figure indicate the number of OTUs comprising 75% of all sequences cloned. Each bin represents an OUT calculated with the furthest-neighbor method a 97% amino-acid similarity cut-off using DOTUR (Schloss and Handelsman, Appl Environ Microbiol 71:1501-1506, 2005). A) *Amitermes* sp. JT5, B) *Gnathamitermes* sp. JT5, C) *Microcerotermes* sp. Cost008, D) *Amitermes* sp. Cost010, E) *Rhyncotermes* sp. Cost004, F) *Nasutitermes* sp. Cost003.